

1 **Supporting Information**

2 For

3 **Phenotyping and Genotyping of Antibiotic-resistance *Escherichia coli* Isolated from**

4 **Beijing River Basin, North China**

5  
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1 **Isolation Procedure of *E. coli*.** Water samples were 10 × fold serial diluted and 0.1 mL of  
2 each dilution was filtered through nitrocellulose filters (0.47 µm pore-size, 47 mm diameter,  
3 Millipore Corporation, America) with the goal of obtaining 30 to 50 colonies per filter. The  
4 filters placed onto *E. coli* chromogenic agar (Chromagar Microbiology, France) and incubated  
5 at 44°C for 24 h. After 24 h of incubation, colonies that turned blue on *E. coli* chromogenic  
6 agar were chosen and streaked onto LB agar (BD, America), and then incubated at 37°C for 24  
7 h. Approximately 40 isolates were collected with dilution method using 6 - 8 disks for each  
8 water sample, and their antibiotic susceptibilities were tested. To avoid the clones, all isolates  
9 were randomly chosen from independent colonies growing on the disks and the number of  
10 selected isolates from each disk was less than 10. The pure cultures were then used to  
11 inoculate 1% tryptone water (Oxoid, UK) and EC broth containing  
12 4-methylumbelliferyl-D-glucuronide (Oxoid, UK) and incubated for 24 h at 37 and 44°C,  
13 respectively. Isolates that produced indole from tryptophan and that were positive for gas  
14 production and fluorescence in EC broth containing 4-methylumbelliferyl-D-glucuronide  
15 were designated as *E. coli* isolates and used for subsequent studies (31, 32).

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**TABLE S1.** PCR Primers used in this study.

	Primers	Target	Sequence (5'- 3')	Annealing temperature(°C)	Amplicon size (bp)	Reference
A	<i>tet</i> (B)/P-FW	<i>tet</i> (B/P)	AAAAC TTATTATATTATAGTG	46	169	(11)
	<i>tet</i> (B)/P-RV		TGGAGTATCAATAATATTCAC			
	<i>tet</i> (M)-FW	<i>tet</i> (M)	ACAGAAAAGCTTATTATATAAC	55	171	(11)
	<i>tet</i> (M)-RV		TGGCGTGTCTATGATGTTTAC			
	<i>tet</i> (O)-FW	<i>tet</i> (O)	ACGGARAGTTTATTGTATACC	60	171	(11)
	<i>tet</i> (O)-RV		TGGCGTATCTATAATGTTGAC			
	<i>tet</i> (Q)-FW	<i>tet</i> (Q)	AGAATCTGCTGTTTGCCAGTG	63	169	(11)
	<i>tet</i> (Q)-RV		CGGAGTGTCAATGATATTGCA			
	<i>tet</i> (S)-FW	<i>tet</i> (S)	GAAAAGCTTACTATACAGTAGC	50	169	(11)
	<i>tet</i> (S)-RV		AGGAGTATCTACAATATTTAC			
	<i>tet</i> (T)-FW	<i>tet</i> (T)	AAGGTTTATTATATAAAAAGTG	46	169	(11)
	<i>tet</i> (T)-RV		AGGTGTATCTATGATATTTAC			
	<i>tet</i> (W)-FW	<i>tet</i> (W)	GAGAGCCTGCTATATGCCAGC	64	168	(11)
<i>tet</i> (W)-RV	GGGCGTATCCACAATGTAAAC					
OTR-FW	<i>Otr</i> (A)	GGCATYCTGGCCACGT	66	212	(11)	
OTR-RV		CCCCGGGTGTCGTASAGG				
B	<i>tet</i> (A)-FW	<i>tet</i> (A)	GCGCGATCTGGTTCACCTCG	61	164	(12)
	<i>tet</i> (A)-RV		AGTCGACAGYRGC GCCGGC			
	<i>tet</i> (B)-FW	<i>tet</i> (B)	TACGTGAATTTATTGCTTCGG	61	206	(12)
	<i>tet</i> (B)-RV		ATACAGCATCCAAGCGCAC			
	<i>tet</i> (C)-FW	<i>tet</i> (C)	GCGGGATATCGTCCATTCCG	68	207	(12)
	<i>tet</i> (C)-RV		GCGTAGAGGATCCACAGGACG			
	<i>tet</i> (D)-FW	<i>tet</i> (D)	GGAAATATCTCCCGGAAGCGG	68	187	(12)
	<i>tet</i> (D)-RV		CACATTGGACAGTGCCAGCAG			
	<i>tet</i> (E)-FW	<i>tet</i> (E)	GTTATTACGGGAGTTTGTTGG	61	199	(12)
	<i>tet</i> (E)-RV		AATACAACACCCACACTACGC			
	<i>tet</i> (G)-FW	<i>tet</i> (G)	CCYGCAAGAGAAGCCAGAAG	68	134	(12)
	<i>tet</i> (G)-RV		CCTTCTCGACCAGGTCGG			
	<i>tet</i> (H)-FW	<i>tet</i> (H)	CAGTGAAAATTCAC TGGAAC	61	185	(12)
	<i>tet</i> (H)-RV		ATCCAAAGTGTGGTTGAGAAT			
	<i>tet</i> (J)-FW	<i>tet</i> (J)	CGAAAACAGACTCGCCAATC	61	184	(12)
	<i>tet</i> (J)-RV		TCCATAATGAGGTGGGGC			
	<i>tet</i> (L)-FW	<i>tet</i> (L)	GATTGGAGTTCTTTGTGGGG	434	55	(35)
	<i>tet</i> (L)-RV		CAATTGCAATACCTGTTCCC			
	<i>tet</i> (Y)-FW	<i>tet</i> (Y)	ATTTGTACCGGCAGAGCAAAC	68	181	(12)
	<i>tet</i> (Y)-RV		GGCGCTGCCGCCATTATGC			
<i>tet</i> (Z)-FW	<i>tet</i> (Z)	CCTTCTCGACCAGGTCGG	61	204	(12)	
<i>tet</i> (Z)-RV		ACCCACAGCGTGTCCGTC				
<i>tet</i> (30)-FW	<i>tet</i> (30)	CATCTTGGTCGAGGTGACTGG	68	134	(12)	
<i>tet</i> (30)-RV		ACGAGCACCCAGCCGAGC				
C	<i>sul</i> (I)-FW	<i>sul</i> (I)	CGCACCGGAAACATCGCTGCAC	55.9	163	(14)
	<i>sul</i> (I)-RV		TGAAGTTCCGCCGCAAGGCTCG			
	<i>sul</i> (II)-FW	<i>sul</i> (II)	TCCGGTGGAGGCCGGTATCTGG	60.8	191	(14)
	<i>sul</i> (II)-RV		CGGGAATGCCATCTGCC TTGAG			
	<i>sul</i> (III)-FW	<i>sul</i> (III)	TCCGTTTCAGCGAATTGGTGACG	60	128	(14)
	<i>sul</i> (III)-RV		TTCGTTACGCCTTACACCAGC			

**CONTIUE TO TABLE S1. PCR Primers used in this study.**

Primers	Target	Sequence (5'- 3')	Annealing temperature(°C)	Amplicon size (bp)	Reference
<i>TEM-FW</i>	<i>TEM</i>	AAAGATGCTGAAGATCA	44	425	(16)
<i>TEM-RV</i>		TTTGGTATGGCTTCATTC			
<i>SHV-FW</i>	<i>SHV</i>	GCGAAAGCCAGCTGTCGGGC	62	304	(16)
<i>SHV-RV</i>		GATTGGCGGCGCTGTTATCGC			
<i>CARB-FW</i>	<i>CARB</i>	CAAGTACTTTYAAAAACAATAGC	46	534	(16)
<i>CARB -RV</i>		GCTGTAATACTCCKAGCAC			

A: Tetracycline resistance genes Encoding Ribosomal Protection Proteins; B: Encoding Tetracycline Efflux Genes; C: encoding sulfonamide resistance gene; D: encoding  $\beta$ -lactamases gene. NA: Not applicable

**TABLE S2.** Detection of Resistance Genes among from Antibiotic-resistance *E.coli* isolates from Beijing River Basin

	Season	Site	Resistance pattern	<i>tet(A)</i>	<i>tet(B)</i>	<i>tet(M)</i>	<i>sul(I)</i>	<i>sul(II)</i>	<i>sul(III)</i>	<i>TEM</i>
01	Summer	A	AMP, TC, SXT	+	-	+	-	+	+	+
02	Summer	A	AMP, TC, SXT, LEV	-	-	-	+	+	-	+
03	Summer	A	AMP, TC, SXT, LEV, CN	+	+	-	+	+	-	+
04	Summer	A	AMP, TC, SXT	+	-	+	+	-	-	+
05	Summer	B	AMP, TC, SXT, LEV	+	+	-	+	-	-	+
06	Summer	B	AMP, TC, SXT	-	+	-	+	+	-	+
07	Summer	B	AMP, TC, SXT	+	-	-	+	+	-	+
08	Summer	B	AMP							+
09	Summer	B	AMP,SXT				+	+	-	+
10	Summer	D	AMP, TC, SXT	+	-	-	-	+	-	+
11	Summer	D	SXT				+	+	-	
12	Summer	E	AMP, TC, SXT	+	+	-	+	+	-	+
13	Summer	E	AMP, TC, SXT, LEV, CN	+	-	-	-	-	+	+
14	Summer	E	AMP,TC,SXT,CN,KZ	-	+	-	+	+	-	+
15	Summer	E	AMP, TC, SXT, LEV, CN	-	+	-	+	+	-	+
16	Summer	E	AMP,TC,SXT,KZ, MA,CFP	-	+	-	+	+	-	+
17	Summer	E	AMP,TC,MA,KZ	-	+	+				+
18	Summer	F	AMP, TC, SXT	+	+	-	+	-	-	+
19	Summer	F	AMP,TC,SXT,KZ,MA	-	-	-	+	+	-	+
20	Summer	F	AMP,TC,SXT,LEV,CN	+	-	-	+	+	-	+
21	Summer	F	AMP,TC,SXT,CN	+	+	-	+	+	-	+
22	Summer	F	AMP,TC,SXT	+	-	-	+	-	-	+
23	Summer	F	AMP,TC,SXT,CN	+	-	-	+	+	-	+
24	Summer	F	AMP,SXT	-	-	-	+	+	-	+
25	Summer	G	AMP,TC.SXT	-	+	+	-	-	-	+

+: present; -: absent

**CONTINUE TO TABLE S2.** Detection of Resistance Genes among from Antibiotic-resistance *E.coli* isolates from Beijing River Basin

	Season	Site	Resistance pattern	<i>tet(A)</i>	<i>tet(B)</i>	<i>tet(M)</i>	<i>sul(I)</i>	<i>sul(II)</i>	<i>sul(III)</i>	<i>TEM</i>
26	Summer	G	AMP,TC,SXT,CN	+	-	-	-	-	+	+
27	Summer	G	AMP,TC,SXT,CN	+	+	-	+	+	-	+
28	Summer	G	TC,SXT	+	-	-	-	+	-	
29	Summer	G	AMP,TC,SXT,CN	+	-	+	-	+	-	+
30	Summer	G	AMP,TC	+	-	-	-	-	-	+
31	summer	G	TC	+	-	-				
32	Summer	G	AMP,TC,SXT	+	-	-	-	-	-	+
33	Summer	H	TC,SXT	+	-	-	+	-	-	
34	Summer	H	AMP,TC,SXT.LEV	-	+	-	+	-	-	+
35	Summer	H	AMP,TC,SXT	+	+	-	-	-	-	+
36	Summer	H	AMP,TC,SXT	-	-	-	-	+	-	+
37	Summer	H	TC,SXT	+		+	+			
38	Summer	H	TC,SXT				+	+		
39	Summer	I	AMP,TC,SXT	+	-	-	+	+	+	+
40	Summer	I	AMP,TC,SXT	+	-	-	+	-	-	+
41	Summer	I	AMP,TC,SXT,LEV,KZ,MA	+	-	-	-	+	+	+
42	Summer	I	AMP,TC,SXT	-	+	-	-	-	-	+
43	Summer	J	TC,SXT	-	+	+	+	-	-	
44	Summer	J	TC,SXT	+	-	+	-	+	-	
45	Summer	K	AMP,TC,SXT,CN	-	-	+	+	+	-	+
46	Summer	K	AMP							-
47	Summer	K	AMP,TC	+	+	-				+
48	Summer	K	AMP,TC	-	-	-				+
49	Summer	K	AMP,TC,SXT,CN	-	+	+	+	+	-	+
50	Summer	K	AMP,TC,SXT,CN	+	-	+	+	+	-	+

+: present; -: absent

**CONTINUE TO TABLE S2.** Detection of Resistance Genes among from Antibiotic-resistance *E.coli* isolates from Beijing River Basin

	Season	Site	Resistance pattern	<i>tet(A)</i>	<i>tet(B)</i>	<i>tet(M)</i>	<i>sul(I)</i>	<i>sul(II)</i>	<i>sul(III)</i>	<i>TEM</i>
51	Summer	K	AMP,TC,SXT,CN	-	-	-	+	+	-	+
52	Winter	A	TC, SXT, LEV	-	-	-	+	-	-	
53	Winter	A	AMP							+
54	Winter	A	TC,SXT	+	-	-	+	+	-	
55	Winter	A	SXT				+	+	+	
56	Winter	A	TC	-	-	-				
57	Winter	A	TC	-	+	-				
58	winter	B	TC	-	+	-				
59	Winter	B	AMP, TC, SXT, LEV	+	-	-	+	+	-	+
60	winter	B	TC	+	-	-				
61	Winter	B	AMP, TC, SXT, LEV	+	-	-	+	-	-	+
62	Winter	B	AMP							+
63	Winter	B	AMP, TC, SXT	-	+	+	+	+	-	+
64	winter	B	TC	+	-	-				
65	winter	B	TC	+	-	-				
66	Winter	B	AMP, TC, SXT,CN	-	+	-	+	+	-	+
67	Winter	C	AMP, TC, SXT, LEV	+	-	-	+	+	-	+
68	Winter	C	AMP,KZ,MA							+
69	Winter	C	AMP,KZ,CFP,MA							+
70	Winter	C	AMP, TC, SXT, LEV	+	-	-	+	+	-	+
71	Winter	C	AMP,SXT				-	+	-	+
72	Winter	C	AMP,SXT,KZ,MA,CN				+	-	-	+
73	Winter	C	AMP,SXT,CN				+	+	+	+
74	Winter	C	TC,SXT	+	-	-	+	+	-	
75	Winter	C	TC,SXT	-	+	-	+	+	-	

+: present; -: absent

**CONTINUE TO TABLE S2.** Detection of Resistance Genes among from Antibiotic-resistance *E.coli* isolates from Beijing River Basin

	Season	Site	Resistance pattern	<i>tet(A)</i>	<i>tet(B)</i>	<i>tet(M)</i>	<i>sul(I)</i>	<i>sul(II)</i>	<i>sul(III)</i>	<i>TEM</i>
76	winter	C	TC	-	+	-				
77	winter	C	TC	-	+	-				
78	winter	C	TC	+	-	-				
79	winter	C	TC	-	+	-				
80	Winter	D	AMP, TC, SXT, LEV, CN	+	-	-	+	+	+	+
81	Winter	D	TC, SXT, LEV, CN	-	+	-	+	+	+	
82	Winter	D	AMP, TC, SXT, LEV, CN	+	+	-	+	+	+	+
83	Winter	D	AMP,KZ							+
84	Winter	D	AMP,KZ,MA							+
85	Winter	D	AMP,TC,KZ	+	-	-				+
86	Winter	D	AMP,TC	-	+	-				+
87	Winter	D	AMP,TC,SXT,KZ,MA,CN	+	+	+	+	+	-	+
88	Winter	D	TC,SXT	+	-	-	+	+	-	
89	winter	D	TC	+	+	-				
90	winter	D	TC	-	+	-				
91	winter	D	TC	+	-	-				
92	Winter	E	AMP, TC, SXT, LEV, CN	+	+	-	+	+	-	+
93	Winter	E	AMP, TC, SXT, LEV, CN	+	-	-	+	+	-	+
94	Winter	E	AMP, TC, SXT, LEV, CN	+	+	-	+	+	-	+
95	Winter	E	TC,SXT,CN	+	-	-	+	+	-	
96	winter	E	TC	+	-	-				
97	winter	E	TC	+	-	-				
98	winter	E	TC	-	+	-				
99	Winter	L	AMP, TC, SXT, LEV	-	+	-	+	+	-	+
100	Winter	L	AMP,TC, SXT, LEV, CN	+	-	-	+	+	-	+

+: present; -: absent



**CONTINUE TO TABLE S2.** Detection of Resistance Genes among from Antibiotic-resistance *E.coli* isolates from Beijing River Basin

	Season	Site	Resistance pattern	<i>tet(A)</i>	<i>tet(B)</i>	<i>tet(M)</i>	<i>sul(I)</i>	<i>sul(II)</i>	<i>sul(III)</i>	<i>TEM</i>
101	Winter	L	AMP, TC, SXT, LEV	+	+	-	+	+	-	+
102	Winter	L	AMP, TC, SXT, LEV	+	-	-	+	+	-	+
103	Winter	L	AMP, TC, SXT, LEV, CN	+	-	-	+	+	-	+
104	Winter	L	AMP							+
105	Winter	L	AMP							-
106	Winter	L	AMP,TC	+	-	+				+
107	Winter	L	AMP,TC,KZ,MA	+	-	+				+
108	Winter	L	AMP,SXT,LEV				+	-	+	+
109	Winter	L	TC,SXT	+	+	-	+	+	+	

+: present; -: absent

AMP: Ampicillin; TC: Tetracycline; SXT: Sulfamethoxazole/trimethoprim; LEV: Levofloxacin; KZ: Cefazolin; MA: Cefamandole; CFP: Cefoperazone; IPM: Imipenem; CN: Gentamicin

**TABLE S3.** Detection of Resistance Genes among Antibiotic-resistance *E.coli* Isolats from animal source and Domestic Wastewater\*. from Sewage Treatment Plant.

	Source	Resistance pattern	<i>tet</i> (A)	<i>tet</i> (B)	<i>tet</i> (M)	<i>sul</i> (I)	<i>sul</i> (II)	<i>sul</i> (III)	<i>TEM</i>
01	A	AMP,TC,SXT	-	+	-	+	-	-	+
02	A	AMP							+
03	A	AMP,TC	-	+	-				+
04	A	AMP,TC	-	+	-				+
05	A	AMP,TC,SXT	+	+	-	+	+	+	+
06	A	AMP,TC,SXT	+	-	-	+	+	-	+
07	A	AMP,LEV,CN							+
08	A	AMP,TC,SXT,LEV	+	-	+	-	-	+	+
09	A	AMP,TC	-	+	-				+
10	A	AMP,TC,SXT,CN	+	-	-	+	+	-	+
11	A	AMP,TC	-	+	-				+
12	A	AMP,TC,SXT	-	+	-	-	-	+	+
13	A	AMP,TC,SXT,CN	+	-	-	-	+	+	+
14	A	AMP,TC,SXT	-	+	-	-	+	+	+
15	A	AMP,TC	-	+	-				+
16	A	AMP,TC,SXT	-	+	-	-	+	+	+
17	A	AMP,TC,SXT,LEV,CN	+	-	-	+	-	+	+
18	A	AMP,TC,SXT	+	-	-	+	+	+	+
19	A	AMP,TC,LEV,CN	-	+	-				+
20	A	AMP,TC,SXT	-	+	-	-	-	-	+
21	A	AMP,TC,SXT	+	+	-	+	-	+	+
22	A	AMP,TC,SXT	+	-	-	+	-	-	+
23	A	AMP,TC,SXT,LEV,CN	+	-	-	-	+	+	+
24	A	AMP,TC	-	+	-				+

+: present; -: absent

**CONTINUE TO TABLE S3.** Detection of Resistance Genes among Antibiotic-resistance *E.coli* Isolats from animal source and Domestic Wastewater\*. from Sewage Treatment Plant.

	Source	Resistance pattern	<i>tet(A)</i>	<i>tet(B)</i>	<i>tet(M)</i>	<i>sul(I)</i>	<i>sul(II)</i>	<i>sul(III)</i>	<i>TEM</i>
25	A	AMP,TC	-	+	-				+
26	A	AMP,TC,SXT	-	-	-	-	+	-	+
27	A	AMP,TC,SXT,LEV	-	+	-	-	-	+	+
28	A	AMP,TC,SXT,LEV	-	-	-	-	+	+	+
29	A	AMP,TC,SXT,LEV,CN	-	-	-	-	-	+	+
30	A	AMP,TC,SXT,LEV,CN	-	-	-	-	+	-	+
31	B	AMP,TC,SXT,LEV,CN	+	-	-	+	+	-	+
32	B	AMP,TC,SXT,LEV	-	+	-	+	-	-	+
33	B	SXT,LEV				+	+	-	
34	B	AMP,TC,SXT,LEV,CN	+	-	-	-	+	+	+
35	B	TC,SXT	-	+	-	-	-	+	
36	B	AMP,TC,SXT,LEV,CN	-	+	-	-	+	-	+
37	B	AMP,TC	+	-	-				+
38	B	AMP,SXT,LEV				-	+	-	+
39	B	AMP,TC,SXT,CN	-	+	-	+	+	-	+
40	B	AMP,TC,SXT,LEV	-	+	-	+	+	-	+
41	B	TC,LEV	-	-	-				
42	B	TC	-	-	-				
43	B	AMP,TC,SXT,LEV	-	+	-	+	-	-	+
44	B	AMP,TC,SXT,LEV,CN	+	-	-	+	+	-	+
45	B	AMP,TC,SXT,LEV	-	-	-	+	+	-	+
46	B	AMP,SXT,LEV				+	-	-	+
47	B	AMP,SXT,LEV				+	-	-	+
48	B	AMP,TC,LEV	-	+	-			-	+

+: present; -: absent

**CONTINUE TO TABLE S3.** Detection of Resistance Genes among Antibiotic-resistance *E.coli* Isolats from animal source and Domestic Wastewater\*. from Sewage Treatment Plant.

	Source	Resistance pattern	<i>tet</i> (A)	<i>tet</i> (B)	<i>tet</i> (M)	<i>sul</i> (I)	<i>sul</i> (II)	<i>sul</i> (III)	<i>TEM</i>
49	B	SXT,LEV				-	+	-	
50	B	SXT,LEV				+	+	-	
51	B	AMP,TC,SXT	-	+	-	-	+	+	+
52	B	TC,SXT,	+	-	-	-	+	-	
53	B	AMP,TC,SXT,LEV,CN	-	+	-	+	+	-	+
54	B	AMP,TC,SXT,	-	-	-	+	+	-	+
55	B	TC,SXT	+	-	-	+	+	-	+
56	B	TC,SXT,LEV	+	-	-	+	+	-	+
57	B	TC	+	-	-				
58	B	AMP,TC,SXT,LEV,CN	-	+	-	+	+	-	+
59	B	AMP,TC,SXT,LEV	+	+	-	+	+	-	+
60	B	AMP,TC,SXT	+	-	-	+	+	-	+
61	B	AMP,TC,SXT,LEV	-	-	-	-	+	-	+
62	B	AMP,TC,SXT,CN	+	+		+	+	-	+
63	B	AMP							+
64	C	TC	+	+	-				
65	C	AMP,TC,SXT,CN	+	-	-	-	+	-	+
66	C	TC	+	-	-				
67	C	AMP,TC,SXT	-	+	-	+	+	+	-
68	C	AMP,TC,SXT	-	-	-	+	+	-	-
69	C	AMP,TC,SXT	+	-	-	+	+	+	-
70	C	AMP,TC,SXT	+	+	-	+	+	+	+
71	C	AMP,TC,SXT,LEV,CN	-	-	-	+	+	-	-
72	C	AMP,TC,SXT	-	-	-	+	-	-	-

+: present; -: absent

**CONTINUE TO TABLE S3.** Detection of Resistance Genes among Antibiotic-resistance *E.coli* Isolats from animal source and Domestic Wastewater\*. from Sewage Treatment Plant.

	Source	Resistance pattern	<i>tet(A)</i>	<i>tet(B)</i>	<i>tet(M)</i>	<i>sul(I)</i>	<i>sul(II)</i>	<i>sul(III)</i>	<i>TEM</i>
73	C	AMP,TC,SXT	+	-	-	+	+	+	-
74	C	AMP,TC,SXT	-	-	-	+	+	+	-
75	C	TC	+	-	-				
76	C	AMP,TC,SXT	+	-	-	+	+	+	-
77	C	AMP,TC,SXT,LEV	-	+	-	+	+	+	-
78	C	AMP,TC,SXT	+	-	-	-	+	-	+
79	C	AMP,TC	+	+	-				+
80	C	TC	+	-	-				
81	C	AMP,TC,SXT,LEV,CN	+	-	-	+	+	+	+
82	C	TC,SXT	+	-	-	+	+	-	
83	C	TC,SXT,LEV	+	-	-	+	+	+	
84	C	AMP,TC,SXT	+	-	-	+	+	-	+
85	C	AMP,TC,SXT,LEV,CN	-	-	-	+	+	-	+
86	C	AMP,TC,SXT	+	-	-	+	-	-	+
87	C	AMP,TC	-	-	-				+
88	C	AMP,TC,SXT,CN	+	-	-	+	+	+	+
89	C	AMP,TC,SXT,LEV	+	-	-	+	+	-	+
90	C	AMP,TC,SXT	+	-	-	+	+	+	+
91	C	AMP,TC,SXT,LEV	-	-	-	+	+	-	-
92	C	TC,SXT	+	+	-	+	+	-	
93	C	AMP,TC,LEV	-	-	-	-	-	-	-
94	C	AMP,TC,SXT	-	-	-	-	+	+	+
95	C	AMP,TC,SXT,LEV	-	-	-	-	+	+	+
96	C	AMP,TC,SXT,LEV	-	-	-	+	+	+	+

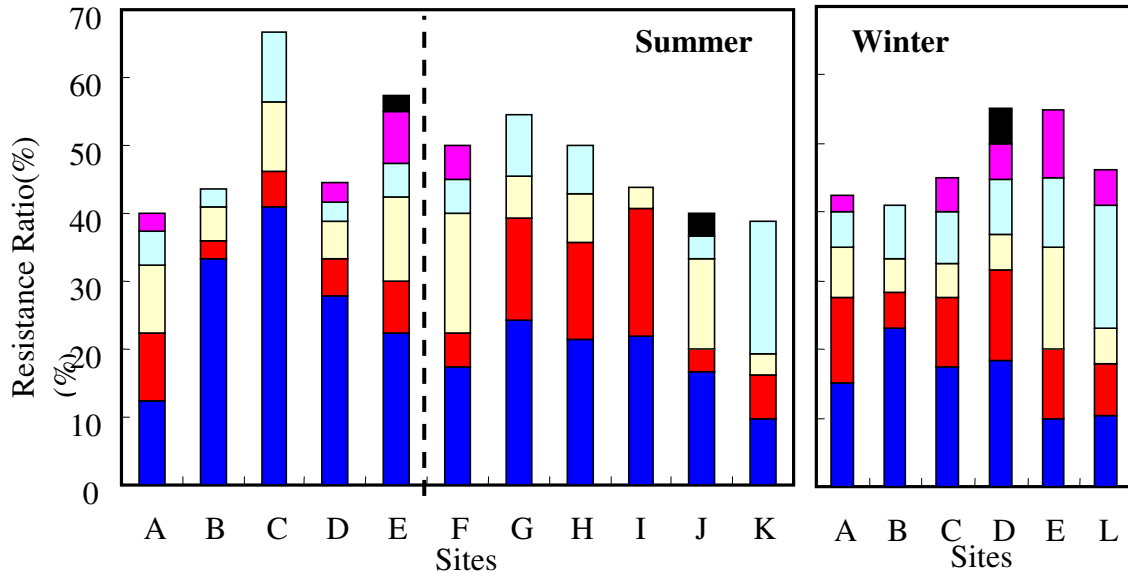
+: present; -: absent

**CONTINUE TO TABLE S3.** Detection of Resistance Genes among Antibiotic-resistance *E.coli* Isolats from animal source and Domestic Wastewater\*. from Sewage Treatment Plant.

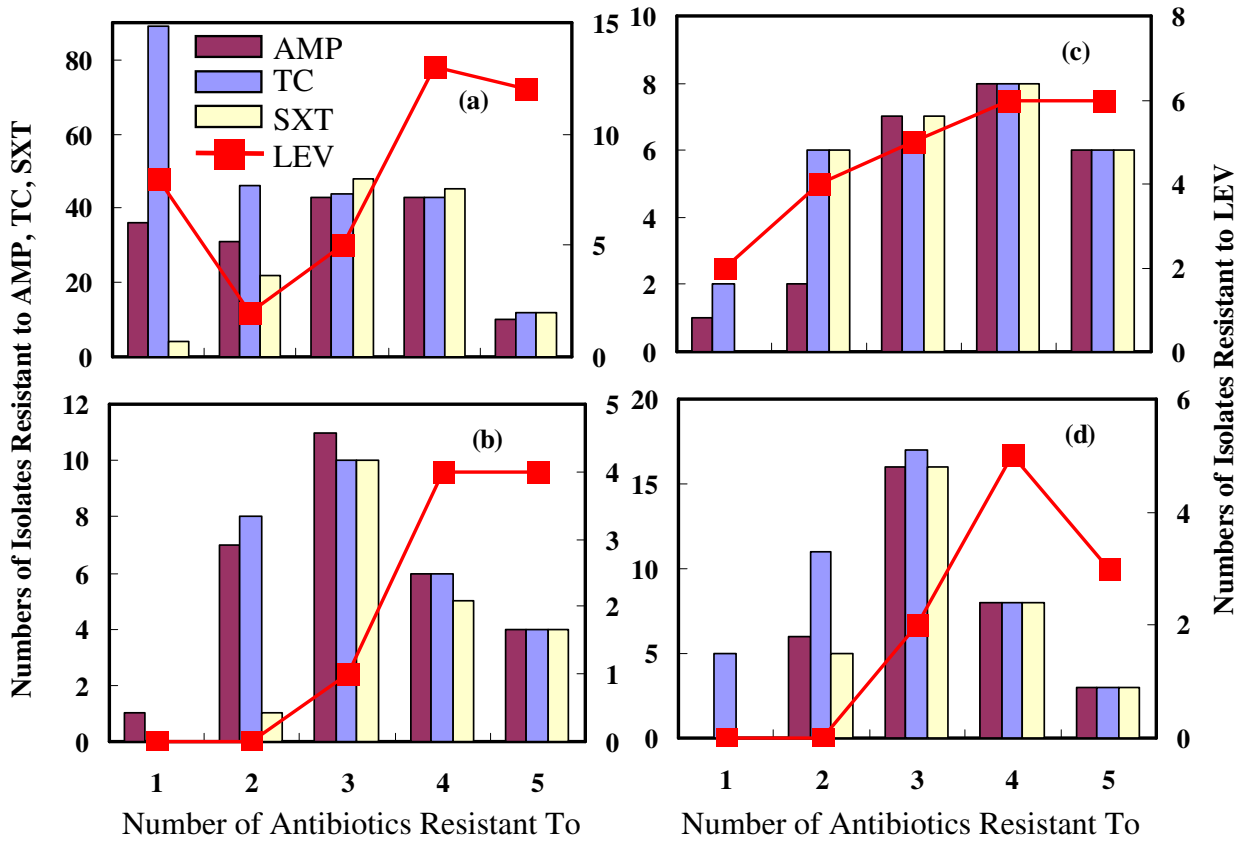
	Source	Resistance pattern	<i>tet(A)</i>	<i>tet(B)</i>	<i>tet(M)</i>	<i>sul(I)</i>	<i>sul(II)</i>	<i>sul(III)</i>	<i>TEM</i>
97	C	TC	+	+	-				
98	C	AMP,TC,SXT	-	-	-	+	+	-	+
99	C	AMP,TC,SXT,CN	+	-	-	+	+	+	+
100	C	TC,SXT	-	-	-	+	+	-	
101	C	AMP,TC	+	+	-				+
102	C	AMP,TC	+	-	-				-
103	C	AMP,TC	+	-	-				+
104	C	AMP,TC	+	-	-				+
105	C	AMP,TC,SXT	-	+	-	+	+	+	-
106	C	TC,SXT	+	-	-	-	+	+	
107	C	AMP,TC,SXT	-	+	-	+	+	+	-

+: present; -: absent

A: swine; B: human; C: chicken; AMP: Ampicillin; TC: Tetracycline; SXT: Sulfamethoxazole/trimethoprim; LEV: Levofloxacin; KZ: Cefazolin; MA: Cefamandole; CFP: Cefoperazone; IPM: Imipenem; CN: Gentamicin; \*Wastewater from Fangzhuang Sewage Treatment Plant in Figure 1.

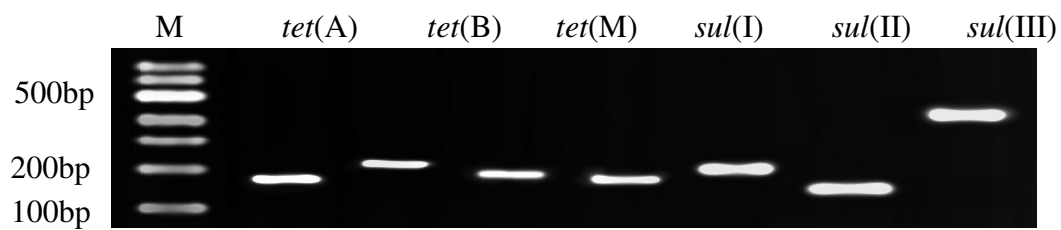


**FIGURE S1.** Profiles of antibiotic-resistance *E.coli* isolated from the Wenyu River Basin. ■ One-drug resistance; ■ two-drug resistance; ■ three-drug resistance; ■ four-drug resistance; ■ five-drug resistance; ■ six-drug resistance.



**FIGURE S2.** Comparison of spectrum of AMP, TC, TMP-SXT, and LEV resistance *E.coli* isolates from Beijing river basin with feedlot and domestic wastewater. AMP: Ampicillin; TC: tetracycline; SXT: sulfamethoxazole-trimethoprim; LEV: levofloxacin. (a) Beijing river basin; 294 antibiotic resistance isolates screened from 624 *E.coli* isolates; (b) swine source; 30 antibiotic resistance isolates screened from 31 *E.coli* isolates; (c) human source from Fangzhuang sewage treatment plant; 36 antibiotic resistance isolates screened from 107 *E.coli* isolates; (d) chicken source; 44 antibiotic resistance isolates screened from 45 *E.coli* isolates.





**FIGURE S3.** Detection of resistance genes coding for tetracycline, sulfonamide and  $\beta$ -Lactam resistance using DNAs from *E.coli* isolated from Wenyu River Basin.